

SKBG-3017US2  
SEQUENCE LISTING

<110> DEEN, KEITH CHARLES  
HURLE, MARK R.  
YOUNG, PETER  
TAN, K.B.

<120> DNA ENCODING TUMOR NECROSIS RELATED RECEPTOR TR7

<130> SKBG-3017US2

<140> TO BE ASSIGNED

<141> 2003-09-15

<150> US 60/041,796

<151> 1997-04-02

<150> US 08/959,382

<151> 1997-10-28

<150> US 09/314,844

<151> 1999-05-19

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<170> PatentIn version 3.2

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Ile Ala Arg Arg Ala Thr Ala Thr Met Ile Ala Gly Ser Leu Leu Leu
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Leu Gly Phe Leu Ser Thr Thr Thr Ala Gln Pro Glu Gln Lys Ala Ser
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 Thr Asn Thr Ser Leu Arg Val Cys Ser Ser Cys Pro Val Gly Thr Phe  
 85 90 95  
 Thr Arg His Glu Asn Gly Ile Glu Lys Cys His Asp Cys Ser Gln Pro  
 100 105 110  
 Cys Pro Trp Pro Met Ile Glu Lys Leu Pro Cys Ala Ala Leu Thr Asp  
 115 120 125  
 Arg Glu Cys Thr Cys Pro Pro Gly Met Phe Gln Ser Asn Ala Thr Cys  
 130 135 140  
 Ala Pro His Thr Val Cys Pro Val Gly Trp Gly Val Arg Lys Lys Gly  
 145 150 155 160  
 Thr Glu Thr Glu Asp Val Arg Cys Lys Gln Cys Ala Arg Gly Thr Phe  
 165 170 175  
 Ser Asp Val Pro Ser Ser Val Met Lys Cys Lys Ala Tyr Thr Asp Cys  
 180 185 190  
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 195 200 205  
 Asn Val Cys Gly Thr Leu Pro Ser Phe Ser Ser Ser Thr Ser Pro Ser  
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 Pro Gly Thr Ala Ile Phe Pro Arg Pro Glu His Met Glu Thr His Glu  
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His His Arg His Ile Leu Lys Leu Leu Pro Ser Met Glu Ala Thr Gly  
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Gly Glu Lys Ser Ser Thr Pro Ile Lys Gly Pro Lys Arg Gly His Pro  
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Arg Gln Asn Leu His Lys His Phe Asp Ile Asn Glu His Leu Pro Trp  
340 345 350

Met Ile Val Leu Phe Leu Leu Leu Val Leu Val Val Ile Val Val Cys  
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Ser Ile Arg Lys Ser Ser Arg Thr Leu Lys Lys Gly Pro Arg Gln Asp  
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Pro Ser Ala Ile Val Glu Lys Ala Gly Leu Lys Lys Ser Met Thr Pro  
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Thr Gln Asn Arg Glu Lys Trp Ile Tyr Tyr Cys Asn Gly His Gly Ile  
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Asp Ile Leu Lys Leu Val Ala Ala Gln Val Gly Ser Gln Trp Lys Asp  
420 425 430

Ile Tyr Gln Phe Leu Cys Asn Ala Ser Glu Arg Glu Val Ala Ala Phe  
435 440 445

Ser Asn Gly Tyr Thr Ala Asp His Glu Arg Ala Tyr Ala Ala Leu Gln  
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His Trp Thr Ile Arg Gly Pro Glu Ala Ser Leu Ala Gln Leu Ile Ser  
465 470 475 480

Ala Leu Arg Gln His Arg Arg Asn Asp Val Val Glu Lys Ile Arg Gly  
485 490 495

Leu Met Glu Asp Thr Thr Gln Leu Glu Thr Asp Lys Leu Ala Leu Pro  
500 505 510

Met Ser Pro Ser Pro Leu Ser Pro Ser Pro Ile Pro Ser Pro Asn Ala  
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Lys Leu Glu Asn Ser Ala Leu Leu Thr Val Glu Pro Ser Pro Gln Asp  
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 tcctgcttgg attccttagc accaccacag ctccagccaga acagaaggcc tcgaatctca 360  
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 cagcaggaac ctatgtctct gagcattgta ccaacacaag cctgcgcgtc tgcagcagtt 480  
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Ile Ala Arg Arg Ala Thr Ala Thr Met Ile Ala Gly Ser Leu Leu Leu  
 20 25 30

Leu Gly Phe Leu Ser Thr Thr Thr Ala Gln Pro Glu Gln Lys Ala Ser  
 35 40 45

Asn Leu Ile Gly Thr Tyr Arg His Val Asp Arg Ala Thr Gly Gln Val  
 50 55 60

Leu Thr Cys Asp Lys Cys Pro Ala Gly Thr Tyr Val Ser Glu His Cys  
 65 70 75 80

Thr Asn Thr Ser Leu Arg Val Cys Ser Ser Cys Pro Val Gly Thr Phe  
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Thr Arg His Glu Asn Gly Ile Glu Lys Cys His Asp Cys Ser Gln Pro  
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Cys Pro Trp Pro Met Ile Glu Lys Leu Pro Cys Ala Ala  
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 gcggcggttg atgcggcgct gggcagaagc agccgccgat tccagctgcc ccgcgcgccc 180  
 cggccacctt gcgagtcctt gggtcagcca tggggacctc tccgagcagc agcaccgccc 240  
 tcggcctcct gcaaccgcat cggccgccga gccacagcca cgatgatcgc gggctccctt 300  
 ctctgtcttg gattccttag caccaccaca gctcagccag aacagaaggc ctcgaatctc 360  
 attggcacat accgccatgt tgaccgtgcc accggccagg tgctaacctg tgacaagtgt 420  
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 20 25 30

Val Asp Arg Ala Thr Gly Gln Val Leu Thr Cys Asp Lys Cys Pro Ala  
 35 40 45

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Gly Thr Tyr Val Ser Glu His Cys Thr Asn Thr Ser Leu Arg Val Cys  
 50 55 60

Gln Gln Cys Pro Val Gly Thr Phe Thr Arg His Glu Asn Gly Ile Glu  
 65 70 75 80

Lys Cys His Asp Cys Ser Gln Pro Cys Pro Trp Pro Met Ile Glu Lys  
 85 90 95

Leu Pro Cys Ala Ser Ala  
 100